

**CSE 421**

**Dynamic Programming**

Shayan Oveis Gharan

# Knapsack Problem

# Knapsack Problem

Given  $n$  objects and a "knapsack."

Item  $i$  weighs  $w_i > 0$  kilograms (an integer) and value  $v_i \geq 0$ .

Knapsack has capacity of  $W$  kilograms.

**Goal:** fill knapsack so as to maximize total value.

**Ex:** OPT is { 3, 4 } with (weight 10) and value 36.

$$W = 11$$

Item	Value	Weight
1	1	2
2	5	3
3	14	4
4	22	6
5	30	8

**Greedy:** repeatedly add item with maximum ratio  $v_i / w_i$ .

**Ex:** { 5, 2 } achieves only value = 35  $\Rightarrow$  greedy not optimal.

# Dynamic Programming: First Attempt

Let  $OPT(i)$  = Max value of subsets of items  $1, \dots, i$  of weight  $\leq W$ .

**Case 1:**  $OPT(i)$  does not select item  $i$

- In this case  $OPT(i) = OPT(i - 1)$

**Case 2:**  $OPT(i)$  selects item  $i$

- In this case, item  $i$  does not immediately imply we have to reject other items
- The problem does not reduce to  $OPT(i - 1)$  because we now want to pack as much value into box of weight  $\leq W - w_i$

**Conclusion:** We need more subproblems, we need to strengthen IH.

# Stronger DP (Strengthening Hypothesis)

Let  $OPT(i, w)$  = Max value subset of items  $1, \dots, i$  of weight  $\leq w$  where  $0 \leq i \leq n$  and  $0 \leq w \leq W$ .

**Case 1:**  $OPT(i, w)$  selects item  $i$

- In this case,  $OPT(i, w) = v_i + OPT(i - 1, w - w_i)$

Take best of the two

**Case 2:**  $OPT(i, w)$  does not select item  $i$

- In this case,  $OPT(i, w) = OPT(i - 1, w)$ .

Therefore,

$$OPT(i, w) = \begin{cases} 0 & \text{If } i = 0 \\ OPT(i - 1, w) & \text{If } w_i > w \\ \max(OPT(i - 1, w), v_i + OPT(i - 1, w - w_i)) & \text{o.w.,} \end{cases}$$

# DP for Knapsack

```
Compute-OPT(i,w)
  if M[i,w] == empty
    if (i==0)
      M[i,w]=0
    else if (wi > w)
      M[i,w]=Comp-OPT(i-1,w)
    else
      M[i,w]= max {Comp-OPT(i-1,w), vi + Comp-OPT(i-1,w-wi) }
  return M[i, w]
```

recursive

# Bottom up DP for Knapsack

```
for w = 0 to W
  M[0, w] = 0
for i = 1 to n
  for w = 1 to W
    if (wi > w)
      M[i, w] = M[i-1, w]
    else
      M[i, w] = max {M[i-1, w], vi + M[i-1, w-wi ]}
return M[n, W]
```

Non-recursive

# DP for Knapsack

←  $W + 1$  →

		0	1	2	3	4	5	6	7	8	9	10	11
$n + 1$	$\phi$	0	0	0	0	0	0	0	0	0	0	0	0
	{1}	0											
	{1, 2}	0											
	{1, 2, 3}	0											
	{1, 2, 3, 4}	0											
	{1, 2, 3, 4, 5}	0											

$W = 11$

```

if ( $w_i > w$ )
     $M[i, w] = M[i-1, w]$  ←
else
     $M[i, w] = \max \{M[i-1, w], v_i + M[i-1, w-w_i]\}$ 
    
```

Item	Value	Weight
1	1	1
2	6	2
3	18	5
4	22	6
5	28	7



# DP for Knapsack

————— W + 1 —————→

		0	1	2	3	4	5	6	7	8	9	10	11
<div style="display: flex; align-items: center;"> <div style="border-left: 1px solid black; border-right: 1px solid black; height: 100px; margin-right: 5px;"></div> <div style="display: flex; flex-direction: column; align-items: center; justify-content: space-around; width: 20px;"> <span>n + 1</span> <span>↓</span> </div> </div>	ϕ	0	0	0	0	0	0	0	0	0	0	0	0
	{ 1 }	0	1	1	1	1	1	1	1	1	1	1	1
	{ 1, 2 }	0											
	{ 1, 2, 3 }	0											
	{ 1, 2, 3, 4 }	0											
	{ 1, 2, 3, 4, 5 }	0											

W = 11

```

if (wi > w)
    M[i, w] = M[i-1, w]
else
    M[i, w] = max {M[i-1, w], vi + M[i-1, w-wi ]}
```

Item	Value	Weight
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# DP for Knapsack

←  $W + 1$  →

		0	1	2	3	4	5	6	7	8	9	10	11
$n + 1$	ϕ	0	0	0	0	0	0	0	0	0	0	0	0
	{1}	0	1	1	1	1	1	1	1	1	1	1	1
	{1, 2}	0	1	6	7								
	{1, 2, 3}	0	1										
	{1, 2, 3, 4}	0	1										
	{1, 2, 3, 4, 5}	0	1										

OPT: { 4, 3 }  
 value = 22 + 18 = 40

$W = 11$

Item	Value	Weight
1	1	1
2	6	2
3	18	5
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5	28	7

```

if (wi > w)
    M[i, w] = M[i-1, w]
else
    M[i, w] = max {M[i-1, w], vi + M[i-1, w-wi ]}
    
```

# DP for Knapsack

←————— W + 1 —————→

		0	1	2	3	4	5	6	7	8	9	10	11
n + 1 ↓	ϕ	0	0	0	0	0	0	0	0	0	0	0	0
	{ 1 }	0	1	1	1	1	1	1	1	1	1	1	1
	{ 1, 2 }	0	1	6	7	7	7	7	7	7	7	7	7
	{ 1, 2, 3 }	0	1	6	7	7	18	19					
	{ 1, 2, 3, 4 }	0	1										
	{ 1, 2, 3, 4, 5 }	0	1										

OPT: { 4, 3 }  
 value = 22 + 18 = 40

W = 11

Item	Value	Weight
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```

if (wi > w)
  M[i, w] = M[i-1, w]
else
  M[i, w] = max {M[i-1, w], vi + M[i-1, w-wi ]}
  
```

# DP for Knapsack

←  $W + 1$  →

		0	1	2	3	4	5	6	7	8	9	10	11
$n + 1$	$\phi$	0	0	0	0	0	0	0	0	0	0	0	0
	{1}	0	1	1	1	1	1	1	1	1	1	1	1
	{1, 2}	0	1	6	7	7	7	7	7	7	7	7	7
	{1, 2, 3}	0	1	6	7	7	18	19	24	25	25	25	25
	{1, 2, 3, 4}	0	1	6	7	7	18	22	24	28	29		
	{1, 2, 3, 4, 5}	0	1										

OPT: { 4, 3 }  
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Item	Value	Weight
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if (wi > w)
    M[i, w] = M[i-1, w]
else
    M[i, w] = max {M[i-1, w], vi + M[i-1, w-wi]}
    
```

# DP for Knapsack

←  $W + 1$  →

		0	1	2	3	4	5	6	7	8	9	10	11
$n + 1$	ϕ	0	0	0	0	0	0	0	0	0	0	0	0
	{1}	0	1	1	1	1	1	1	1	1	1	1	1
	{1,2}	0	1	6	7	7	7	7	7	7	7	7	7
	{1,2,3}	0	1	6	7	7	18	19	24	25	25	25	25
	{1,2,3,4}	0	1	6	7	7	18	22	24	28	29	29	40
	{1,2,3,4,5}	0	1	6	7	7	18	22	28	29	34	34	40

OPT: { 4, 3 }  
 value = 22 + 18 = 40

$W = 11$

Item	Value	Weight
1	1	1
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3	18	5
4	22	6
5	28	7

```

if (wi > w)
    M[i, w] = M[i-1, w]
else
    M[i, w] = max {M[i-1, w], vi + M[i-1, w-wi ]}
    
```

# Knapsack Problem: Running Time

Running time:  $\Theta(n \cdot W)$

- Not polynomial in input size!
- "Pseudo-polynomial."
- Decision version of Knapsack is NP-complete.

Knapsack approximation algorithm:

There exists a polynomial algorithm that produces a feasible solution that has value within 0.01% of optimum in time  $\text{Poly}(n, \log W)$ .

# DP Ideas so far

- You may have to define an ordering to decrease #subproblems
- $\text{OPT}(i,w)$  is exactly the predicate of induction
- You may have to strengthen DP, equivalently the induction, i.e., you may have to carry more information to find the Optimum.
- This means that sometimes we may have to use two dimensional or three dimensional induction

# RNA Secondary Structure

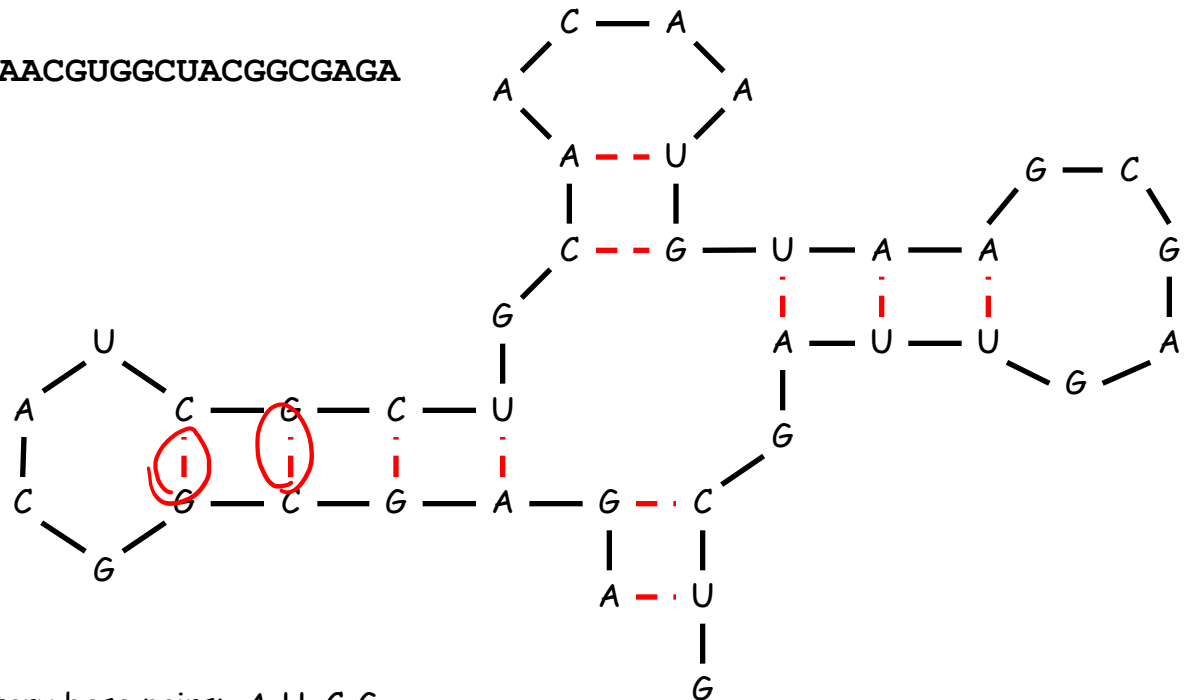


# RNA Secondary Structure

**RNA:** A String  $B = b_1b_2\dots b_n$  over alphabet  $\{ A, C, G, U \}$ .

**Secondary structure.** RNA is single-stranded so it tends to loop back and form **base pairs** with itself. This structure is essential for understanding behavior of molecule.

**Ex:** GUCGAUUGAGCGAAUGUAACAACGUGGCUACGGCGAGA



complementary base pairs: A-U, C-G

# RNA Secondary Structure (Formal)

**Secondary structure.** A set of pairs  $S = \{ (b_i, b_j) \}$  that satisfy:

[Watson-Crick.]

- $S$  is a *matching* and
- each pair in  $S$  is a Watson-Crick pair: A-U, U-A, C-G, or G-C.

[No sharp turns.]: The ends of each pair are separated by at least 4 intervening bases. If  $(b_i, b_j) \in S$ , then  $i < j - 4$ .

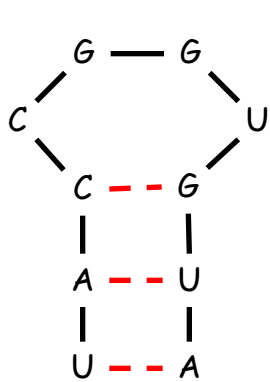
[Non-crossing.] If  $(b_i, b_j)$  and  $(b_k, b_l)$  are two pairs in  $S$ , then we cannot have  $i < k < j < l$ .

**Free energy:** Usual hypothesis is that an RNA molecule will maximize total free energy.

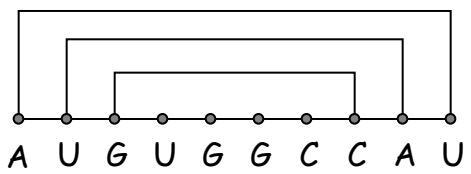
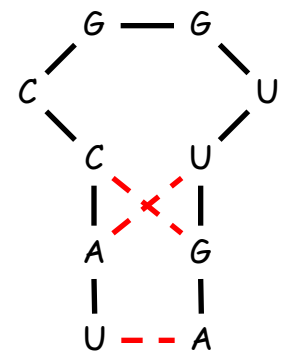
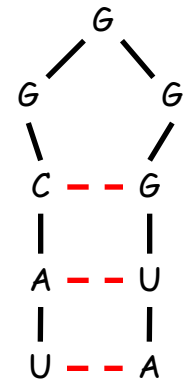
↙  
approximate by number of base pairs

**Goal:** Given an RNA molecule  $B = b_1b_2\dots b_n$ , find a secondary structure  $S$  that maximizes the number of base pairs.

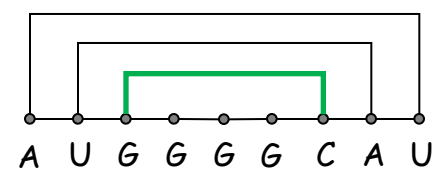
# Secondary Structure (Examples)



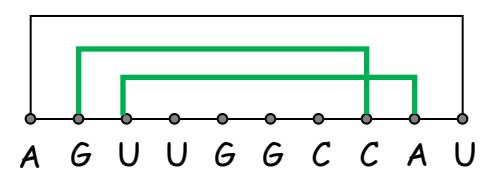
↖  
base pair



ok



~~sharp turn~~



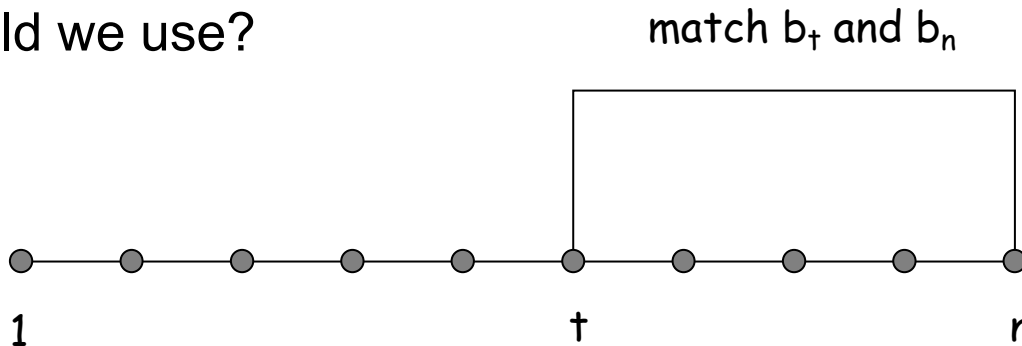
~~crossing~~

# DP: First Attempt

**First attempt.** Let  $OPT(n)$  = maximum number of base pairs in a secondary structure of the substring  $b_1b_2\dots b_n$ .

Suppose  $b_n$  is matched with  $b_t$  in  $OPT(n)$ .

What IH should we use?



**Difficulty:** This naturally reduces to two subproblems

- Finding secondary structure in  $b_1, \dots, b_{t-1}$ , i.e.,  $OPT(t-1)$
- Finding secondary structure in  $b_{t+1}, \dots, b_{n-1}$ , ???

# DP: Second Attempt

**Definition:**  $OPT(i, j)$  = maximum number of base pairs in a secondary structure of the substring  $b_i, b_{i+1}, \dots, b_j$

The most important part of a correct DP; It fixes IH

**Case 1:** If  $j - i \leq 4$ .

- $OPT(i, j) = 0$  by no-sharp turns condition.

**Case 2:** Base  $b_j$  is not involved in a pair.

- $OPT(i, j) = OPT(i, j-1)$

**Case 3:** Base  $b_j$  pairs with  $b_t$  for some  $i \leq t < j - 4$

- non-crossing constraint **decouples** resulting sub-problems
- $OPT(i, j) = \max_{t: b_i \text{ pairs with } b_t} \{ 1 + OPT(i, t - 1) + OPT(t + 1, j - 1) \}$

# Recursive Code

Let  $M[i,j]$ =empty for all  $i,j$ .

```
Compute-OPT(i,j){
  if (j-i <= 4)
    return 0;
  if (M[i,j] is empty)
    M[i,j]=Compute-OPT(i,j-1)
  for t=i to j-5 do
    if ( $b_t, b_j$  is in {A-U, U-A, C-G, G-C})
      M[i,j]=max(M[i,j], 1+Compute-OPT(i,t-1) +
        Compute-OPT(t+1,j-1))
  return M[j]
}
```

Does this code terminate?  
What are we inducting on?

# Formal Induction

Let  $OPT(i, j)$  = maximum number of base pairs in a secondary structure of the substring  $b_i, b_{i+1}, \dots, b_j$

**Base Case:**  $OPT(i, j) = 0$  for all  $i, j$  where  $|j - i| \leq 4$ .

**IH:** For some  $\ell \geq 4$ , Suppose we have computed  $OPT(i, j)$  for all  $i, j$  where  $|i - j| \leq \ell$ .

**IS:** Goal: We find  $OPT(i, j)$  for all  $i, j$  where  $|i - j| = \ell + 1$ . Fix  $i, j$  such that  $|i - j| = \ell + 1$ .

**Case 1:** Base  $b_j$  is not involved in a pair.

- $OPT(i, j) = OPT(i, j-1)$  [this we know by IH since  $|i - (j - 1)| = \ell$ ]

**Case 2:** Base  $b_j$  pairs with  $b_t$  for some  $i \leq t < j - 4$

- $OPT(i, j) = \max_{t: b_i \text{ pairs with } b_t} \{ 1 + OPT(i, t - 1) + OPT(t + 1, j - 1) \}$

We know by IH since difference  $\leq \ell$

# Bottom-up DP

```
for k = 1, 2, ..., n-1
  for i = 1, 2, ..., n-1
    j = i + k
    if (j-i <= 4)
      M[i,j]=0;
    else
      M[i,j]=M[i,j-1]
      for t=i to j-5 do
        if ( $b_t, b_j$  is in {A-U, U-A, C-G, G-C})
          M[i,j]=max(M[i,j], 1+ M[i,t-1] + M[t+1,j-1])

return M[1, n]
}
```

4	0	0	0	↗
3	0	0	↗	↗
2	0	↗	↗	↗
1	↗	↗	↗	↗
	6	7	8	9

j

Running Time:  $O(n^3)$



# Lesson

We may not always induct on  $i$  or  $w$  to get to smaller subproblems.

We may have to induct on  $|i - j|$  or  $i + j$  when we are dealing with more complex problems, e.g., intervals